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(54) Title: SYSTEMS AND METHODS FOR MICROARRAY DATA ANALYSIS

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\mathsf{EM\_ESTIMATE}\;(\mu_{\!\scriptscriptstyle 1},\,...,\,\mu_{\!\scriptscriptstyle K},\,\Sigma_{\!\scriptscriptstyle 1},\,...,\,\Sigma_{\!\scriptscriptstyle K},\,\tau_{\!\scriptscriptstyle 1},\,...,\,\tau_{\!\scriptscriptstyle K},\,\mathit{A'})
                           FOR EACH ROW R OF A' WITH MISSING VALUES
                           FOR i = 1, ..., K
                           USE EM AND \textit{N}(\mu_{j_1},\Sigma_{j}) TO ESTIMATE THE
                      MISSING VALUES IN R.
   R_i \leftarrow R WITH MISSING VALUES REPLACED BY ESTIMATES.
   R' \leftarrow \text{WEIGHTEDAVERAGE}(R_1, ..., R_k).
  REPLACE R IN A' BY R'.
  RETURN A'.
  K_ESTIMATE(K, A)
                         /* FIRST PART: INITIALIZATION */
  B ← ROWS OF A WITHOUT MISSING VALUES.
  \begin{array}{c} \mu_1, \, ..., \, \mu_K, \, \Sigma_1, \, ..., \, \Sigma_K, \, \tau_1, \, ..., \, \tau_K \leftarrow \\ \text{GAUSSIAN MIXTURE CLUSTERING OF } \textit{B}. \end{array}
\begin{array}{l} \textit{A'} \leftarrow \text{EM\_ESTIMATE} \; (\mu_1, \, ..., \, \mu_K, \, \Sigma_1, \, ..., \, \Sigma_K, \, \tau_1, \, ..., \, \tau_K, \, \textit{A}). \\ \textit{/*} \; \text{SECOND PART: ITERATION */} \end{array}
 REPEAT
                          \mu_1, ..., \mu_K, \Sigma_1, ..., \Sigma_K, \tau_1, ..., \tau_K
       GAUSSIAN MIXTURE CLUSTERING OF A'.
 A' \leftarrow \mathsf{EM\_ESTIMATE} \ (\mu_1, \, ..., \, \mu_K, \, \Sigma_1, \, ..., \, \Sigma_K \, \tau_1, \, ..., \, \tau_K, \, A').
 ) UNTIL CONVERGENCE
GMCimpute(S, A)
                         FOR K = 1, 2, ..., S
                        A_{K} \leftarrow K_{ESTIMATE}(K, A).
RETURN (A1+ A2 + ... + AS) / S.
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(57) Abstract: Clustering is routinely applied in the exploratory analysis of microarray data. Missing entries arise from blemishes on the microarrays. The present invention provides a new method, and computer program and/or computer product thereof to impute missing values. The method involves the steps of clustering microarray data by partitioning the data into a select number of clusters, wherein each data point is iteratively moved from one cluster to another, until two consecutive iterations have resulted in the same partition pattern; obtaining a select number of estimates of the data in the clusters by probabilistic interference; and averaging the select number of estimates to obtain missing values in the microarray data. The method is superior to other imputation models as measured by root mean squared errors.

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